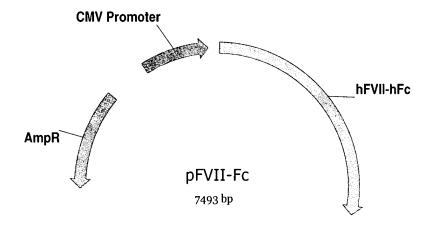
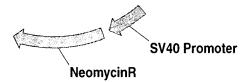
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FIGURE 1 - Plasmid vector pFVII-Fc according to example 1

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FIGURE 2

SEQ ID NO:1 (The amino acid sequence of native human coagulation Factor VII):

5

Ala-Asn-Ala-Phe-Leu-GLA-GLA-Leu-Arg-Pro-Gly-Ser-Leu-GLA-Arg-GLA-Cys-Lys5 10 15

GLA-GLA-Gln-Cys-Ser-Phe-GLA-GLA-Ala-Arg-GLA-Ile-Phe-Lys-Asp-Ala-GLA-Arg10 20 25 30 35

Thr-Lys-Leu-Phe-Trp-Ile-Ser-Tyr-Ser-Asp-Gly-Asp-Gln-Cys-Ala-Ser-Ser-Pro-40 45 50

Cys-Gln-Asn-Gly-Gly-Ser-Cys-Lys-Asp-Gln-Leu-Gln-Ser-Tyr-Ile-Cys-Phe-Cys55 60 65 70

Leu-Pro-Ala-Phe-Glu-Gly-Arg-Asn-Cys-Glu-Thr-His-Lys-Asp-Asp-Gln-Leu-Ile-75 80 85 90

20

35

Cys-Val-Asn-Glu-Asn-Gly-Gly-Cys-Glu-Gln-Tyr-Cys-Ser-Asp-His-Thr-Gly-Thr-95 100 105

Lys-Arg-Ser-Cys-Arg-Cys-His-Glu-Gly-Tyr-Ser-Leu-Leu-Ala-Asp-Gly-Val-Ser-25 110 115 120 125

Cys-Thr-Pro-Thr-Val-Glu-Tyr-Pro-Cys-Gly-Lys-Ile-Pro-Ile-Leu-Glu-Lys-Arg130 135 140

30 Asn-Ala-Ser-Lys-Pro-Gln-Gly-Arg-Ile-Val-Gly-Gly-Lys-Val-Cys-Pro-Lys-Gly-145 150 155 160

Glu-Cys-Pro-Trp-Gln-Val-Leu-Leu-Leu-Val-Asn-Gly-Ala-Gln-Leu-Cys-Gly-Gly-165 170 175 180

Thr-Leu-Ile-Asn-Thr-Ile-Trp-Val-Val-Ser-Ala-Ala-His-Cys-Phe-Asp-Lys-Ile185 190 195

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SEQ ID NO:2 (DNA primer 1 for preparation of hFVII-hFc):

5'- GCTAGCCACCATGGTCTCCCAGGCCCTCAG -3' (SEQ ID NO:2)

5 SEQ ID NO:3 (DNA primer 2 for preparation of hFVII-hFc):

5'- CGAGCCCCATTTCCCGGATCCGCAGAGCCCAAATCTTGT -3' (SEQ ID NO:3)

SEQ ID NO:4 (DNA primer 3 for preparation of hFVII-hFc):

5'- CGAGCCCCATTTCCCGGATCCGCAGAGCCCAAATCTTGT -3' (SEQ ID NO:4)

10

SEQ ID NO:5 (DNA primer 4 for preparation of hFVII-hFc):

5'- TTGCCGGCCGTCGCACTCATTTA -3' (SEQ ID NO:5)

SEQ ID NO:6 (The amino acid sequence of native human coagulation Factor VII with alternative spliced propeptide (underlined) conjugated to Fc domain of IgG1, native human coagulation Factor VII underlined.:

MVSQALRLLCLLLGLQGCLAAGGVAKASGGETRDMPWKPGPHRVFVTQEEAHGVLHRRRRANAF
LEELRPGSLERECKEEQCSFEEAREIFKDAERTKLFWISYSDGDQCASSPCQNGGSCKDQLQSYICF

20 CLPAFEGRNCETHKDDQLICVNENGGCEQYCSDHTGTKRSCRCHEGYSLLADGVSCTPTVEYPCGKI
PILEKRNASKPQGRIVGGKVCPKGECPWQVLLLVNGAQLCGGTLINTIWVVSAAHCFDKIKNWRNLIA
VLGEHDLSEHDGDEQSRRVAQVIIPSTYVPGTTNHDIALLRLHQPVVLTDHVVPLCLPERTFSERTLAF
VRFSLVSGWGQLLDRGATALELMVLNVPRLMTQDCLQQSRKVGDSPNITEYMFCAGYSDGSKDSCK
GDSGGPHATHYRGTWYLTGIVSWGQGCATVGHFGVYTRVSQYIEWLQKLMRSEPRPGVLLRAPFP

25 GSAEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYV
DGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREP
QVYTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDK
SRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK

30

SEQ ID NO:7 (Amino acid sequence of Fc domain from IgG1):

EPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGV EVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVY

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TLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSR WQQGNVFSCSVMHEALHNHYTQKSLSLSPGK

5

SEQ ID NO:8 (The amino acid sequence of native human coagulation Factor VII conjugated to Fc domain of IgG1, native human coagulation Factor VII underlined, X refers to GLA residues):

10 ANAFLXXLRPGSLXRXCKXXQCSFXXARXIFKDAXRTKLFWISYSDGDQCASSPCQNGGSCKDQLQS
YICFCLPAFEGRNCETHKDDQLICVNENGGCEQYCSDHTGTKRSCRCHEGYSLLADGVSCTPTVEYP
CGKIPILEKRNASKPQGRIVGGKVCPKGECPWQVLLLVNGAQLCGGTLINTIWVVSAAHCFDKIKNWR
NLIAVLGEHDLSEHDGDEQSRRVAQVIIPSTYVPGTTNHDIALLRLHQPVVLTDHVVPLCLPERTFSER
TLAFVRFSLVSGWGQLLDRGATALELMVLNVPRLMTQDCLQQSRKVGDSPNITEYMFCAGYSDGSK
15 DSCKGDSGGPHATHYRGTWYLTGIVSWGQGCATVGHFGVYTRVSQYIEWLQKLMRSEPRPGVLLR
APFPGSAEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFN
WYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQP
REPQVYTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLT

VDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK

20

SEQ ID NO:9 (cDNA sequence encoding the amino acid sequence of native human coagulation Factor VII with alternative spliced propeptide conjugated to Fc domain of IgG1:

atggtctcccaggccctcaggctcctctgccttctgctttgggcttcagggctgcctggctgcaggcggggtcgctaaggcctcaggagagaaa 25 aacgcgttcctggaggagctgcggccgggctccctggagagggagtqcaaggaggagtqccattqctccttcgaggaggagcccqqqagatcttc aaggacgcggagaggacgaagctgttctggatttcttacagtgatggggaccagtgtgcctcaagtccatgccagaatgggggctcctgcaag gaccagctccagtcctatatctgcttctgcctccctgccttcgagggccggaactgtgagacgcacaaggatgaccagctgatctgtgtgaacg agaacggcggctgtgagcagtactgcagtgaccacacgggcaccaagcgctcctgtcggtgccacgaggggtactctctqctqqcaqacqq ggtgtcctgcacacccacagttgaatatccatgtggaaaaatacctattctagaaaaaagaaatgccagcaaaccccaaggccgaattgtgg 30 ggggcaaggtgtgccccaaaggggagtgtccatggcaggtcctgttgttggtgaatggagctcagttgtgtggggggaccctgatcaacacca tctgggtggtctccgcggcccactgtttcgacaaaatcaagaactggaggaacctgatcgcggtgctgggcgagcacgacctcagcgagcac ccgcctgcaccagcccgtggtcctcactgaccatgtggtgcccctctgcccgaacggacgttctctgagaggacgctggccttcgtgcgctteteattggteageggetggggeeagetgetggaeegtggegeeaeggeeetggageteatggtgeteaaegtgeeeeggetgatgaeeeagg 35 actgcctgcagcagtcacggaaggtgggagactccccaaatatcacggagtacatgttctgtgccggctactcqqatggcagcaaggactcct gcgcgggggacagtggaggccacatgccaccactaccggggcacqtqqtacctgacggcatcqtcagctgaggccagaggtgcac accgtgggccactttggggtgtacaccagggtctcccagtacatcgagtggctgcaaaagctcatgcgctcagagccacgcccaggagtcctc ctgcgagcccatttcccggatccgcagagcccaaatcttgtgacaaaactcacacatgcccaccgtgcccagcacctgaactcctgggggg accytcagtcttcctcttcccccaaaacccaaggacaccctcatqatctcccggacccttqaqtcacatqqtqqtqqtqqacqtqaqccac 40 gaagaccctgaggtcaagttcaactggtacgtggacggcgtggaggtgcataatqccaagacaaagccgcgggaggagcaqtacaacaq cacqtaccqtgtggtcaqcqtcctcaccqtcctgcaccaggactggctgaatggcaaggagtacaagtgcaaggtctccaacaaagccctcc cagccccatcgagaaaaccatctccaaagccaaaggcagccccgagaaccacaggtgtacaccctqcccccatcccqqqatqaqctq 

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aacaactacaagaccacgcctcccgtgctggactccgacggctccttcttcctctacagcaagctcaccgtggacaagagcaggtggcagca ggggaacgtcttctcatgctccgtgatgcatgaggctctgcacaaccactacacgcagaagagcctctccctgtctccgggtaaatga

SEQ ID NO:10 (cDNA sequence encoding the vector comprising the cDNA sequence encoding the amino acid sequence of native human coagulation Factor VII with alternative spliced propeptide conjugated to Fc domain of IgG1:

gctagccaccatggtctcccaggccctcaggctcctctgccttctgctttgggcttcagggctgcctggctgcaggcggggtcgctaaggcctcag 10 gaggagaaacacgggacatgccgtggaagccggggcctcacagagtcttcgtaacccaggaggaagcccacggcgtcctgcaccggcg ccggcgcccaacgcgttcctggaggagctgcggccgggctccctggagagggagtgcaaggaggagtgctccttcgaggaggccc gggagatcttcaaggacgcggagaggacgaagctgttctggatttcttacagtgatgggaccagtgtgcctcaagtccatgccaqaatggg gctcctgcaaggaccagctccagtcctatatctgcttctgcctccctgccttcgagggccggaactgtgagacgcacaaggatgaccagctgat ctgtgtgaacgagaacggctgtgagcagtactgcagtgaccacacgggcaccaagcgctcctgtcggtgccacgaggggtactctctgc 15 ccgaattgtggggggcaaggtgtgccccaaaggggagtgtccatggcaggtcctgttgttggtgaatggagctcagttgtgtggggggaccctg cagegageaegaeggggatgageagageeggegggtggegeaggteateateeceageaegtaegteeegggeaeeaeeaaecaegae 20 cettegtgegetteteattggteageggetgggggeeagetggtggecegtgggeceaggeeetggageteatggtgeteaacgtgeeeegget gcaaggactcctgcgcgggggacagtggaggcccacatgccaccactaccggggcacgtggtacctgacgggcatcgtcagctggggcc agggctgcgcaaccgtgggccactttggggtgtacaccagggtctcccagtacatcgagtggctgcaaaagctcatgcgctcagagccacgc ccaggagtcctcctgcgagccccatttcccggatccgcagagcccaaatcttgtgacaaaactcacacatgcccaccgtgcccagcacctgallowers and the compact of the c25 actectggggggaccgtcagtcttcctcttcccccaaaacccaaggacacctcatgatctcccggacccctgaggtcacatgcgtggtggt gacgtgagccacgaagaccctgaggtcaagttcaactggtacgtggacggcgtggaggtgcataatgccaagacaaagccgcgggagga gcagtacaacagcacgtaccgtgtggtcagcgtcctcaccgtcctgcaccaggactggctgaatggcaaggagtacaagtgcaaggtctcca acaaagccctcccagcccccatcgagaaaaccatctccaaagccaaagggcagccccgagaaccacaggtgtacaccctgcccccatcc 30 ggcagccggagaacaactacaagaccacgcctcccgtgctggactccgacggctccttcttcctctacagcaagctcaccgtggacaagag caggtggcagcaggggaacgtcttctcatgctccgtgatgcatgaggctctgcacaaccactacacgcagaagagcctctccctgtctccgggt aaatgaaagggcgaattctgcagatatccagcacagtggcggccgctcgagtctagagggcccgtttaaacccgctgatcagcctcgactgtg cettetagttgccagecatetgttgtttgcccteccccgtgccttccttgaccctggaaggtgccacteccactgtcctttcctaataaaatgaggaa 35 catgctggggatgcggtgggctctatggcttctgaggcggaaagaaccagctggggctctagggggtatccccacgcgccctgtagcggcgc ccacgttcgccggctttccccgtcaagctctaaatcgggggctccctttagggttccgatttagtgctttacggcacctcgaccccaaaaaacttga ttagggtgatggttcacgtagtgggccatcgccctgatagacggtttttcgccctttgacgttggagtccacgttctttaatagtggactcttqttccaa actggaacaacactcaaccctatctcggtctattcttttgatttataagggattttgccgatttcggcctattggttaaaaaatgagctgatttaacaaa 40 agtecegecetaacteegeceatecegeceetaacteegeceagtteegeceatteteegeceatggetgactaatttttttatttatgeagagg ccgaggccgcctctgcctctgagctattccagaagtagtgaggaggcttttttggaggcctaggcttttgcaaaaagctcccgggagcttgtatat ccattttcggatctgatcaagagacaggatgaggatcgtttcgcatgattgaacaagatggattgcacgcaggttctccggccgcttgggtggagatgattgcacgcaggttctccggccgcttgggttggagatgattgcacgcaggttctccggccgcttgggttggagatgattgcacgcaggttctccggccgcttgggttggagatgattgcacgcaggttctccggccgcttgggttggagatgattgcacgcaggattgcacgcaggttctccggccgcttgggttggagatgattgcacgcaggattgcacgattg45 aggetatteggetatgaetgggeacaacagacaateggetgetetgatgeegeegtgtteeggetgteagegeaggggegeeeggttetttttgte aagaccgacctgtccggtgccctgaatgaactgcaggacgaggcagcgcggctatcgtggctaggccacgacgggggttccttgcgcagctgt gctcgacgttgtcactgaagcgggaagggactggctgctattgggcgaagtgccggggcaggatctcctgtcatctcaccttgctcctgccgag aaagtatecateatggetgatgeaatgeggetgeataegettgateeggetaeetgeeeattegaceaeeaagegaaacategeategaq cgagcacgtactcggatggaagccggtcttgtcgatcaggatgatctggacgaagagcatcaggggctcgcgccagccgaactgttcgccag 50 gctcaaggcgcgcatgcccgacggcgaggatctcgtcgtgacccatggcgatgcctgcttgccgaatatcatggtggaaaatggccgcttttct ggattcatcgactgtggccggctgggtgtggcggaccgctatcaggacatagcgttggctacccgtgatattgctgaagagcttggcggcaat gggetgacegetteetegtgetttaeggtategeegeteeegattegeagegeategeettetategeettettgaegagttettetgageggaetet ggggttcgaaatgaccgaccaagcgacgcccaacctgccatcacgagatttcgattccaccgccgccttctatgaaaggttgggcttcggaat cgttttccgggacgccggctggatgatcctccagcgggggatctcatgctggagttcttcgcccaccccaacttgtttattgcagcttataatggtt 55

acaaataaagcaatagcatcacaaatttcacaaataaagcattttttcactgcattctagttgtggtttgtccaaactcatcaatgtatcttatcatgtc

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tgtataccgtcgacctctagctagagcttggcgtaatcatggtcatagctgtttcctgtgtgaaattgttatccgctcacaattccacacaacatacga5 aacatgtgagcaaaaggccagcaaaaggccaggaaccgtaaaaaggccgcgttgctggcgtttttccatagqctccqccccctgacgagc atcacaaaaatcgacgctcaagtcagaggtggcgaaacccgacaggactataaagataccaggcgtttccccctggaagctccctcgtgcg ctctcctgttccgaccctgccgcttaccggatacctgtccgcctttctcccttcgggaagcgtggcgctttctcatagctcacgctgtaggtatctcagt10 teggtgtaggtcgttcgctccaagctgggctgtgtgcacgaaccccccgttcagcccgaccgctgcgccttatccggtaactatcgtcttgagtccaacccggtaagacacgacttatcgccactggcagcagcactggtaacaggattagcagagcgaggtatgtaggcggtgctacagagttctt gaagtggtggcctaactacggctacactagaagaacagtatttggtatctgcgctctgctgaagccagttaccttcggaaaaagagttggtagct atcttttctacggggtctgacgctcagtggaacgaaaactcacgttaagggattttggtcatgagattatcaaaaaggatcttcacctagatcctttt 15 aaattaaaaatgaagttttaaatcaatctaaagtatatatgagtaaacttggtctgacagttaccaatgcttaatcagtgaggcacctatctcagcg atetgtctatttegttcatccatagttgcctgactccccgtcgtgtagataactacgatacgggagggcttaccatctgqccccagtqctgcaatgat cgtggtgtcacgctcgttcgtttggtatggcttcattcagctccggttcccaacgatcaaggcgagttacatgatcccccatgttgtgcaaaaaaagcg 20 catecgtaagatgettttetgtgaetggtgagtaeteaaceaagteattetgagaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeeggegteaatagtgtatgeggegaeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgetettgeeegagttgeegagttgeeegagttgeeegagttgeeegagttgeeegagttgeeegagttgeeegagttgeeegagtgeeegagttgeeacgggataataccgcgccacatagcagaactttaaaagtgctcatcattggaaaacgttcttcggggcgaaaactctcaaggatcttaccgctg ttgagatccagttcgatgtaacccactcgtgcacccaactgatcttcagcatcttttactttcaccagcgtttctgggtgagcaaaaacaggaaggc aaaatgccgcaaaaaagggaataagggcgacacggaaatgttgaatactcatactcttccttttttcaatattattqaaqcatttatcaqqqttattqt 25 ctcatgagcggatacatatttgaatgtatttagaaaaataaacaaataggggttccgcgcacatttccccgaaaagtgccacctgacgtcgacg aggtcgctgagtagtgcgcgagcaaaatttaagctacaacaaggcaaggcttgaccgacaattgcatgaagaatctgcttagggttaggcgttt tgcgctgcttcgcgatgtacgggccagatatacgcgttgacattgattattgactagttattaatagtaatcaattacggggtcattagttcatagccc atatatggagttccgcgttacataacttacggtaaatggcccgcctggctgaccgcccaacgacccccgcccattgacgtcaataatgacgtat 30 gttcccatagtaacgccaatagggactttccattgacgtcaatgggtggagtatttacggtaaactgcccacttggcagtacatcaaggtatcata tgccaagtacgccccctattgacgtcaatgacggtaaatggcccgcctggcattatgcccagtacatgaccttatgggactttcctacttggcagt acatctacgtattagtcatcgctattaccatggtgatgcggttttggcagtacatcaatgggcgtggatagcggtttgactcacggggatttccaagt ctccaccccattgacgtcaatgggagtttgttttggcaccaaaatcaacgggactttccaaaatgtcgtaacaactccqccccattgacgcaaat gggcggtaggcgtgtacggtgggaggtctatataagcagagctctctggctaactagagaacccactgcttactggcttatcgaaattaatacg 35 actcactatagggagacccaagctg

SEQ ID NO:11 (The amino acid sequence of native human coagulation Factor VII with propertide (underlined) conjugated to Fc domain of IgG1, native human coagulation Factor VII underlined. Construct is made according to example 1):

40 MVSQALRLCLLLGLQGCLAAVFVTQEEAHGVLHRRRRANAFLEELRPGSLERECKEEQCSFEEAR EIFKDAERTKLFWISYSDGDQCASSPCQNGGSCKDQLQSYICFCLPAFEGRNCETHKDDQLICVNEN GGCEQYCSDHTGTKRSCRCHEGYSLLADGVSCTPTVEYPCGKIPILEKRNASKPQGRIVGGKVCPKG ECPWQVLLLVNGAQLCGGTLINTIWVVSAAHCFDKIKNWRNLIAVLGEHDLSEHDGDEQSRRVAQVII PSTYVPGTTNHDIALLRLHQPVVLTDHVVPLCLPERTFSERTLAFVRFSLVSGWGQLLDRGATALELM VLNVPRLMTQDCLQQSRKVGDSPNITEYMFCAGYSDGSKDSCKGDSGGPHATHYRGTWYLTGIVS WGQGCATVGHFGVYTRVSQYIEWLQKLMRSEPRPGVLLRAPFPGSAEPKSCDKTHTCPPCPAPELL GGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRV VSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQVSLTCLVK GFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHY TQKSLSLSPGK

SEQ ID NO:12 (cDNA sequence encoding the amino acid sequence of native human coagulation Factor VII with propertide conjugated to Fc domain of IgG1, cDNA construct is made according to example 1):

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#### 8/9

atggtctcccaggccctcaggctcctctgccttctgctttgggcttcagggctgcctggctgcagtcttcgtaacccaggaggaagcccacggcgtcctgcaccggcgccgcgcccaacgcgttcctggaggagctgcggcccgggctccctggaggaggagtgcaaggaggagcagtgctccttcgaggaggcccgggagatcttcaaggacgcggagaggacgaagctgttctggatttcttacagtgatggggaccagtgtgcctcaagtccatgccagaatgggggctcctgcaaggaccagctccagtcctatatctgcttctgcctccctgccttcgagggccggaactgtgagacgcacaaggat gaccagctgatctgtgtgaacgagaacggcggctgtgagcagtactgcagtgaccacacgggcaccaagcgctcctgtcggtgccacgagg ggtactctctgctggcagacggggtgtcctgcacacccacagttgaatatccatgtggaaaaatacctattctagaaaaaagaaatgccagca aaccccaaggccgaattgtggggggcaaggtgtgccccaaaggggagtgtccatggcaggtcctgttgttggtgaatggagctcagttgtgtg gggggaccctgatcaacaccatctgggtggtctccgcggcccactgtttcgacaaaatcaagaactggaggaacctgatcgcggtgctgggc gaggacgctggccttcgtgcgcttctcattggtcagcggctggggccagctgggaccgtgggcgccacggccctggagctcatggtgctcaac gtgccccggctgatgacccaggactgcctgcagcagtcacggaaggtgggagactccccaaatatcacggagtacatgttctgtgccggcta cagctggggccagggctgcgcaaccgtgggccactttggggtgtacaccagggtctcccagtacatcgagtggctgcaaaagctcatgcgctcagage caa agc caa agage caa agc caa agage caa agc cccagcacctgaactcctggggggaccgtcagtcttcctcttcccccaaaacccaaggacaccctcatgatctcccggacccctgaggtcacat gcgtggtggacgtgagccacgaagaccctgaggtcaagttcaactggtacgtggacggcgtggaggtgcataatgccaagacaaagccacgacgtggacgtgaggtgcataatgccaagacaaagccacgacgtggacgtggacgtgaggtgcataatgccaagacaaagccacgacgtgaggtgacgtgaggtgacgtgaggtgaagttcaactggtacgtgaggtgacgtgaggtgaagttcaactggtacgtgaggtgaaggtgaagttcaactggtacgtgaggtgaaggaagggcgggaggagcagtacaacagcacgtaccgtgtggtcagcgtcctcaccgtcctgcaccaggactggctgaatggcaaggagtacaagtgc a agg to tecaa caa agg cecte ceage cece categagaa aa ceatete caa agg ceaa agg geage ceegagaa ceae agg t g ta cae cette caa agg geage ceegagaa ceae agg t g ta cae cette caa agg geage ceegagaa ceae agg t g ta cae cette ceaa agg geage ceegagaa ceae agg t g ta cae cette ceaa agg geage ceegagaa ceae agg t g ta cae cette ceaa agg geage ceegagaa ceae agg t g ta cae cette ceaa agg ceae agg geage ceegagaa ceae agg t g ta cae cette ceaa agg ceae agg g cae ceegagaa ceae agg t g ta cae cee ceae agg t g ta cae ceae agg t g t ac cae ceae agg t g ta cae ceae agg t g t ac cae ceae agggacaagagcaggtggcagcaggggaacgtcttctcatgctccgtgatgcatgaggctctgcacaaccactacacgcagaagagcctctccctgtctccgggtaaatga

SEQ ID NO:13 (cDNA sequence encoding the vector comprising the cDNA sequence encoding the amino acid sequence of native human coagulation Factor VII with propertide conjugated to Fc domain of IgG1, cDNA construct is made according to example 1):

30 gctagccaccatggtctcccaggccctcaggctcctctgccttctgctttgggcttcagggctgcctggctgcagtcttcgtaacccaggaggaagc ccacggcgtcctgcaccggcgccgcgccaacgcgttcctggaggagctgcgggctccctggagagggagtgcaaggaggagc agtgctccttcgaggaggcccgggagatcttcaaggacgcggagaggacgaagctgttctggatttcttacagtgatggggaccagtgtgcctc aagtccatgccagaatgggggctcctgcaaggaccagctccagtcctatatctgcttctgcctccctgccttcgagggccggaactgtgagacg cacaaggatgaccagctgatctgtgtgaacgagaacggcggctgtgagcagtactgcagtgaccacacgggcaccaagcgctcctgtcggt 35 gccacgaggggtactctctgctggcagacggggtgtcctgcacacccacagttgaatatccatgtggaaaaatacctattctagaaaaaaagaaatgccagcaaaccccaaggccgaattgtggggggcaaggtgtccccaaaggggagtgtccatggcaggtcctgttgttggtgaatggagct cagt tg tg tg tg tg tg tg tg ta acaccat ct tg tg tg tg tc tc cg cg g cccact g tt tc gacaaa at caa gaact ga gg aacct g at cg cg transfer to the transfer transfer to the transfer transfer to the transfer transfer transfer to the transfer transfer40 cgttctctgagaggacgctgggccttcgtgcgcttctcattggtcagcggctggggccagctgctggaccgtggggccacggccctggagctcatg gccggctactcggatggcagcaaggactcctgcggggggacagtggaggcccacatgccaccactaccggggcacgtggtacctgacggg catcg t cag ctg gg gc cag gg ctg ca accg tgg gc cactt tgg gg tg tacaccag gg tctcccag tacatcg ag tg gc tacaccg tgg gc cactt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg caa aa gc tacatcg tgg gc cactt tgg gg tg tacaccag gg tctcccag tacatcg ag tg cac tgg gg ctg cact tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tctcccag tacatcg ag tgg ctg cac tt tgg gg tg tacaccag gg tgg ctg cac tt tgg gg tgg tacaccag gg tgg ctg cac tt tgg gg tgg tacaccag gg tgg tacaccag gg tgg tgg tacaccag gg tgg tgg tacaccag gg tgg tgg tacaccag gg tgg tacaccag gg tgg tgg tgg tacaccag gg tgg tgg tacaccag gg tacaccag gg tgg tacaccag gg tacaccag gg tgg tacaccag gg tgg tacaccag gg tacaccag gcatgcgctcagagcccaggagtcctcctgcgagccccatttcccggatccgcagagcccaaatcttgtgacaaaactcacacatgcc45 caccgtgcccagcacctgaactcctggggggaccgtcagtcttcctcttcccccaaaaacccaaggacaccctcatgatctcccggacccctgaggtcacatgcgtggtggtggacgtgagccacgaagaccctgaggtcaagttcaactggtacgtggacggcgtggaggtgcataatgccaagtcacatgcgaggtgcataatgccaagtcacatgcgaggtggaggtgcataatgccaagtcacatgcgtggaggtggaggtgcataatgccaagtcacatgcgtggaggtggaggtgcataatgccaagtcacatgcgaggtgaaggtgaaggtgcataatgccaagtcacatgcgaggtgaaggtgaaggtgcataatgccaagtcacatgcgaggtgaaggaaggtgaaggtgaaggtgaaggtgaaggtgaaggtgaaggtgaaggtgaaggtgaagaagacaaagccgcgggaggagcagtacaacagcacgtaccgtgtggtcagcgtcctcaccgtcctgcaccaggactggctgaatggcaagga gtacaagtgcaaggtctccaacaaagccctcccagcccccatcgagaaaaccatctccaaagccaaagggcagccccgagaaccacag 50 gtggagtgggagagcaatgggcagccggagaacaactacaagaccacgcctcccgtgctggactccgacggctccttcttcctctacagca agctcaccgtggacaagagcaggtggcagcaggggaacgtcttctcatgctccgtgatgcatgaggctctgcacaaccactacacgcagaa gagcctctccctgtctccgggtaaatgaaagggcgaattctgcagatatccagcacagtggcggccgctcgagtctagagggcccgtttaaac 55 gattgggaagacaatagcaggcatgctggggatgcggtgggctctatggcttctgaggcggaaagaaccagctggggctctagggggtatcc

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